Modeling Frequency and Severity of Claims with the Generalized Cluster-Weighted Model

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Abstract

In this paper, we propose a generalized cluster-weighted model (GCWM) that allows for modeling non-Gaussian distribution of the continuous covariates. Additionally, our zero-inflated GCWM allows for modeling zero-inflated cluster weighted distribution of claims that is more suitable in the insurance applications. We describe an expectation-optimization (EM) algorithm for parameter estimation in GCWM. Cluster-weighted models are considered as a flexible family of mixture models for fitting the joint distribution of a random vector composed of a response variable and a set of continues and discrete covariates. However, these models have a few limitations when it comes to the insurance applications and may provide suboptimal results. A simulation study showed that the GCWM a performs well for different settings in contrast to the existing mixture-based approaches. A real data set based on French auto-mobile policies is used to illustrate the application of the proposed model.

KEY WORDS: finite mixture models, GLM, GCWM a, CWM, ratemaking, automobile claims.

JEL CLASSIFICATION: C02, C40, C60.

1   Introduction

Predictive modeling gained a lot of attention in the past decade in the area of actuarial science, risk management, and insurance in general. While the term predictive modeling has been used in many other areas, in context of insurance, it is referred to as a process of leveraging statistics in estimating the insurance cost (see Frees and Meyers, 2014). Various predictive models are used in the area of actuarial science with generalized

linear models (GLMs) being the most popular tools actively integrated in pricing, reserving, and underwriting of property and casualty insurance. The most recent extensions of the GLM models proposed by Garrido et al. (2016) and Shi et al. (2015) allow for relaxing the assumption of independence between number and size of claims. Several GLM extensions based on copulas have been considered (e.g., Frees et al., 2016; Krämer et al., 2013; Czado et al., 2012; Frees and Wang, 2006).

Ingrassia and Minotti (2015) proposed a cluster-weighted models (CWMs) as a flexible family of mixture models for fitting the joint distribution of a random vector composed of a response variable and a set of mixed-type covariates with the assumption that continues covariates come from Gaussian distribution. In this paper paper, we consider two extensions of CWM model to allow modeling of non-Gaussian continues covariates and a zero-inflated Poisson (ZIP) claims distribution. We define our proposed model as cluster-weighted transformed model or GCWM a which is more suitable for the insurance applications. The CWM models with Gaussian assumptions have been proposed by Gershenfeld (1997), Gershenfeld and Metois (1999), and Gershenfeld (1999) in a context of media technology. Some extensions of this class of models have been considered by Punzo and Ingrassia (2014), Ingrassia and Punzo (2014), and Ingrassia and Vittadini (2014).

There is a growing interest in modeling insurance losses using mixture models. Several recent mixture models of univariate insurance data have been developed, including work by Lee and Lin (2010), Verbelen et al. (2015), and Miljkovic and Grün (2016). A finite mixture of bivariate Poisson regression models with an application to insurance ratemaking was studied by Bermúdez and Karlis (2012). The authors used the EM algorithm to determine the number of components in the mixture. Another Poisson mixture model for count data was considered by Brown and Buckley (2015) with application in managing a Group Life insurance portfolio. Motivated by the idea of mixture modeling, we believe that the extension of the univariate mixture modeling can be applied to mixture modeling of regressions including the GLMs, where losses are modeled as a function of several covariates.

This paper is organized as follows. Section 2 presents the proposed model for mixture of GLMs. Section 3 applies the proposed model on a real data of French automobile claims. An extensive simulation study is discussed in Section 4. Conclusion is provided in Section 5.

2   Proposed Model

2.1   Background

Let $(\boldsymbol{X}', Y)'$ be the pair of a vector of covariates $\boldsymbol{X}$ and a response variable Y. Assume this set is defined on some space Ω that takes values in appropriate Euclidian subspace. Now, assume that there exists G partitions of

Ω, denoted as $\Omega_1, \ldots, \Omega_G$. Gershenfeld (1997) characterized the Cluster weighted models as a finite mixture of GLMs hence, the joint distribution $f(\boldsymbol{x}, y)$ of $(\boldsymbol{X}', Y)'$ is expressed as follows

$$f(\boldsymbol{x}, y) = \sum_{j=1}^{G} \tau_j q(y|\boldsymbol{x}; \Omega_j) p(\boldsymbol{x}; \Omega_j). \tag{2.1}$$

The pair $q(y|\boldsymbol{x}; \Omega_j)$ and $p(\boldsymbol{x}; \Omega_j)$ are conditional and marginal distributions of $(\boldsymbol{X}', Y)'$ respectively, while τ_j represents the weight of the jth component such that $\sum_{j=1}^{G} \tau_j = 1$, $\tau_j > 0$. Ingrassia and Minotti (2015) proposed a flexible family of mixture models for fitting the joint distribution of a random vector $(\boldsymbol{X}', Y)'$ by splitting the covariates into continues and discrete as $\boldsymbol{X} = (\boldsymbol{V}', \boldsymbol{W}')'$. This assumption of independence between continues and discrete covariates allows us to multiply their corresponding marginal distributions. Thus, for this setting the model in (2.1) is reformulated as follows

$$f(\boldsymbol{x}, y; \boldsymbol{\Phi}) = \sum_{j=1}^{G} \tau_j q(y|\boldsymbol{x}; \boldsymbol{\vartheta_j}) p(\boldsymbol{x}; \boldsymbol{\theta_j}) = \sum_{j=1}^{G} \tau_j q(y|\boldsymbol{x}; \boldsymbol{\vartheta_j}) p(\boldsymbol{v}; \boldsymbol{\theta_j^\star}) p(\boldsymbol{w}; \boldsymbol{\theta_j^{\star\star}}) \tag{2.2}$$

where $\boldsymbol{v}$ and $\boldsymbol{w}$ are the vectors of continues and discrete covariates respectively, the $q(y|\boldsymbol{x}; \boldsymbol{\vartheta_j})$ is a conditional density of $Y|\boldsymbol{x}$, with parameter vector $\boldsymbol{\vartheta_j}$, the $p(\boldsymbol{v}; \boldsymbol{\theta_j^\star})$ is the marginal distribution of $\boldsymbol{v}$ with parameter vector $\boldsymbol{\theta_j^\star}$. the $p(\boldsymbol{w}; \boldsymbol{\theta_j^{\star\star}})$ is the marginal distribution of $\boldsymbol{w}$ with parameter vector $\boldsymbol{\theta_j^{\star\star}}$. Finally $\boldsymbol{\Phi} := (\boldsymbol{\theta^\star}, \boldsymbol{\theta^{\star\star}}, \boldsymbol{\tau}, \boldsymbol{\vartheta})$ includes all model parameters. In addition, the conditional distribution $q(y|\boldsymbol{x}; \boldsymbol{\vartheta_j})$ is assumed to belong to an exponential family of distributions and as such can be modeled in the framework of GLMs. Here, the marginal distribution of continues covariates is assumed to be of Gaussian type. Unfortunately, this last assumption is too strong for use in insurance related applications specifically in rate-making. To relax it, we develop the Generalized cluster-weighted mdoel (GCWM) that allows for non-Gaussian covariates as discussed in the next section.

2.2 Generalized cluster-weighted model (GCWM)

We proceed to extend (2.2) by splitting the continues covariates further as $\boldsymbol{V} := (\boldsymbol{U}', \boldsymbol{T}')'$, where $\boldsymbol{U}$ is a set of non-Gaussian covariates, and $\boldsymbol{T}$ a set of Gaussian covariates. Thus (2.2) is now recovered as

$$f(\boldsymbol{x}, y; \boldsymbol{\Phi}) = \sum_{j=1}^{G} \tau_j q(y|\boldsymbol{x}; \boldsymbol{\vartheta_j}) p(\boldsymbol{t}; \boldsymbol{\theta_j^\star}) p(\boldsymbol{w}; \boldsymbol{\theta_j^{\star\star}}) p(\boldsymbol{u}; \boldsymbol{\theta_j^{\star\star\star}}) \tag{2.3}$$

which we refer to as generalized cluster-weighted model (GCWM). Here $p(\boldsymbol{t}; \boldsymbol{\theta_j^\star})$ denotes the marginal density of Gaussian covariates, with parameter vector $\boldsymbol{\theta^\star}$, and $p(\boldsymbol{u}; \boldsymbol{\theta_j^{\star\star\star}})$ as the marginal density of the non-Gaussian covariates with parameter vector $\boldsymbol{\theta_j^{\star\star\star}}$.

As it is relevant to the actuarial application in this paper, we focus on the multivariate log-normal distri-

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bution for non-Gaussian covariates. This however does not reduce the generality of our approach. Now, with our log-normal assumption for $p(\boldsymbol{u}; \boldsymbol{\theta_j^{\star\star\star}})$, we have that $\boldsymbol{u}$ is defined on $\mathbb{R}_+$ with parameter vector $\boldsymbol{\theta_j^{\star\star\star}}$, and having probability density function as

$$p\left(\boldsymbol{u}; \boldsymbol{\theta_j^{\star\star\star}} := (\boldsymbol{\mu_j^{\star\star\star}}, \boldsymbol{\Sigma_j^{\star\star\star}})\right) = \frac{1}{(\prod_{i=1}^N u_i)|\boldsymbol{\Sigma_j^{\star\star\star}}|(2\pi)^{\frac{p}{2}}} \exp\left[-\frac{1}{2}(\ln \boldsymbol{u} - \boldsymbol{\mu_j^{\star\star\star}})' \boldsymbol{\Sigma_j^{\star\star\star-1}}(\ln \boldsymbol{u} - \boldsymbol{\mu_j^{\star\star\star}})\right].$$

(2.4)

The derivation of the equation above can be found in the Appendix 7.1.

2.3   Zero-inflated Poisson

For the zero-inflated Poisson model (ZIP) see Lambert (1992) we can split the conditional density $p(y|\boldsymbol{x}, \boldsymbol{\vartheta_j})$ into zero and non-zero densities. The response y variable when $y = 0$ are distributed with density $q(0|\boldsymbol{x}, \boldsymbol{\vartheta_j})$. The response y values when $y > 0$ are distributed with density $q(y > 0|\boldsymbol{x}, \boldsymbol{\vartheta_j})$. Given the conditional density now defined for the ZIP model, (2.3) can be re-written as follows

$$f(\boldsymbol{x}, y; \Phi) = \sum_{j=1}^G \tau_j \left[q(0|\boldsymbol{x}; \boldsymbol{\vartheta_j}) + q(y > 0|\boldsymbol{x}; \boldsymbol{\vartheta_j}))\right] p(\boldsymbol{t}; \boldsymbol{\theta_j^{\star}}) p(\boldsymbol{w}; \boldsymbol{\theta_j^{\star\star}}) p(\boldsymbol{u}; \boldsymbol{\theta_j^{\star\star\star}}).$$

(2.5)

Let $\tilde{\boldsymbol{x}} := [\boldsymbol{1}, \boldsymbol{x}]$, where $\tilde{\boldsymbol{x}}$ is a matrix of covariates with the addition of a placeholder for the intercept in the GLM. We denote the Poisson conditional density as $q^P(y|\boldsymbol{x}; \boldsymbol{\beta_j})$, where $y \in \{0, 1, \dots\}$, and $\boldsymbol{\beta_j}$ is the row coefficient vector. Here, the link function will be modelled with log-link for the GLM:

$$\lambda_j = e^{\tilde{\boldsymbol{x}}\boldsymbol{\beta_j'}}, \qquad\qquad q^P(y|x; \lambda_j) = e^{-\lambda_j}\frac{\lambda_j{}^y}{y!}.$$

Next, we introduce a Bernoulli process for the conditional density. We denote the density as $q^B(y|\boldsymbol{x}; \bar{\boldsymbol{\beta}}_j)$, where $\mathbb{1}(\cdot)$ is the indicator function and $\bar{\boldsymbol{\beta}}_j$ is the coefficient vector. Here, the GLM will be modeled with the associated logit link function

$$\psi_j = \frac{e^{\tilde{\boldsymbol{x}}\bar{\boldsymbol{\beta}}_j{}'}}{1 + e^{\tilde{\boldsymbol{x}}\bar{\boldsymbol{\beta}}_j{}'}}, \qquad\qquad q^B(y|\boldsymbol{x}; \bar{\boldsymbol{\beta}}_j) = \begin{cases} \psi_j, & \mathbb{1}(y) = 0 \\ 1 - \psi_j, & \mathbb{1}(y) = 1 \end{cases}$$

Now, given a combination of two preceding models, we introduce the ZIP process in which zero counts come from two random variables. One comes from Bernoulli random variable which generates structural zeros, and the other comes from the Poisson random variable. The coefficients $\boldsymbol{\vartheta_j} := \{\boldsymbol{\beta_j}, \bar{\boldsymbol{\beta}}_j\}$ correspond to the two above introduced conditional densities where the coefficients are estimated using a generalized linear model as

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in Lambert (1992). The components of ZIP conditional density $q(y|\boldsymbol{x}; \boldsymbol{\vartheta_j})$ are

$$q(0|x; \boldsymbol{\vartheta_j}) = \psi_j + (1 - \psi_j)e^{-\lambda_j} \qquad \text{and} \qquad q(y > 0|x; \boldsymbol{\vartheta_j}) = (1 - \psi_j)e^{-\lambda_j}\frac{(\lambda_j)^y}{y!}.$$

Also, the link functions to consider are log-link for the Poisson process and logit link for the Bernoulli

$$\psi_j = \frac{e^{\tilde{\boldsymbol{x}}\bar{\boldsymbol{\beta}}_j{}'}}{1 + e^{\tilde{\boldsymbol{x}}\bar{\boldsymbol{\beta}}_j{}'}} \qquad \text{and} \qquad \lambda_j = e^{\tilde{\boldsymbol{x}}\beta_j'}.$$

Let paramater ψ_j denote the probability that the zero comes from the Bernoulli distribution of jth component, and the parameter λ_j characterizes the jth Poisson distribution. This allows for a more nuanced approach to handling the inflation of zeros similarly as in Bermúdez and Karlis (2012).

3   Introducing Bernoulli-Poisson Partitioning

The single component ZIP model assumes that the inflated zeros emanate from both a Bernoulli and Poisson random variables while the non-zeros are assumed to come exclusively from the Poisson random variable. However, recent research extends the single component ZIP models to mixture models for heterogeneous count data with excess zeros (see (Bermúdez and Karlis, 2012)). In mixtures of ZIPs, zeros are assumed to come from multiple different Binomial and Poisson random variables. Difficulties are apparaent during the maximization step of the EM when means of covariates are very close together (see Lim et al. (2014)) . However, misclassification error can be reduced using parsimonious models for the indepedent variables as in McNicholas et al. (2010).

In this work, we propose a new method to rectify this problem and partition the dataset using Bernoulli and Poisson GCWMs. Furthermore, we construct a new zero inflated GCWM (ZI-GCWM) using the previously generated Bernoulli and Poisson GCWMs. We show that the Bernoulli and Poisson GCWM accurately estimate the initialization of the EM algorithm for the zero inflated GCWM model. The work of Lambert (1992) specifies that the MLE estimates for coefficients provide an excellent guess allowing EM to converge quickly for ZIPs. The partitioning method consists of two seperate EM algorithms. The first EM algorithm is for generating the GCWM models, while the second EM is for optimizing the ZI-GCWM. Recall $(\boldsymbol{X}', Y)'$ to be a vector defined on some sample space Ω. As discussed, this sample space is partitioned into G non-overlapping sets such that their union constitutes this sample space ie. $\Omega = \bigcup_{i=1}^{G} \Omega_i$. However, contingent on a model choice each particular set Ω_i may take a different shape. Specifically, if we introduce the Bernoulli model in a generalized form for conditional density (see Ingrassia and Minotti (2015) for specific cases), we have the sample space

Ω^B and joint pdf f^B to be

$$\Omega^B = \bigcup_{l=1}^{G} \Omega_l^B \qquad \text{and} \qquad f^B(\boldsymbol{x}, y; \Phi) = \sum_{l=1}^{G} \tau_l q^B(y|\boldsymbol{x}; \bar{\boldsymbol{\beta}}_{\boldsymbol{l}}) p(\boldsymbol{t}; \boldsymbol{\theta}_{\boldsymbol{l}}^{\star}) p(\boldsymbol{w}; \boldsymbol{\theta}_{\boldsymbol{l}}^{\star\star}) p(\boldsymbol{u}; \boldsymbol{\theta}_{\boldsymbol{l}}^{\star\star\star}).$$

Similarly if we introduce a Poisson model in a generalized form the sample space Ω^P and joint pdf f^P become

$$\Omega^P = \bigcup_{j=1}^{M} \Omega_j^P \qquad \text{and} \qquad f^P(\boldsymbol{x}, y; \Phi) = \sum_{j=1}^{M} \tau_j q^P(y|\boldsymbol{x}; \boldsymbol{\beta}_{\boldsymbol{j}}) p(\boldsymbol{t}; \boldsymbol{\theta}_{\boldsymbol{j}}^{\star}) p(\boldsymbol{w}; \boldsymbol{\theta}_{\boldsymbol{j}}^{\star\star}) p(\boldsymbol{u}; \boldsymbol{\theta}_{\boldsymbol{j}}^{\star\star\star}).$$

Where this sample space is partitioned up to M non-overlapping sets. Now, construct a new partitioning of a sample space Ω such that

$$\Omega = \Omega^Z = \bigcup_{l \in \{1...G\}, j \in \{1...M\}} \Omega_{l,j}^Z := \bigcup_{l \in \{1...G\}, j \in \{1...M\}} \Omega_l^B \cap \Omega_j^P =: \bigcup_{k \in \{1...K\}} \Omega_k^Z$$

, where K can range up to $M \times G$ unique partitions. Therefore the new conditional density is now result of a model in which each component is captured by the conditional probability density function that is of mixture of particular Bernoulli and particular Poisson

$$q_k^Z(y|\boldsymbol{x}; \bar{\boldsymbol{\beta}}_k, \boldsymbol{\beta}_k) := q^B(y|\boldsymbol{x}; \bar{\boldsymbol{\beta}}_k) + (1 - q^B(y|\boldsymbol{x}; \bar{\boldsymbol{\beta}}_k)) q^P(y|\boldsymbol{x}; \boldsymbol{\beta}_k). \tag{3.1}$$

$$= q(0|\boldsymbol{x}; \boldsymbol{\vartheta}_{\boldsymbol{k}}) + q(y > 0|\boldsymbol{x}; \boldsymbol{\vartheta}_{\boldsymbol{k}}) \tag{3.2}$$

The expectation-maximization (EM) algorithm (see Dempster et al. (1977)) is then used to estimate this new mixture of up to $M \times G$ specific GCWMs. The initialization parameters for the second EM algorithm are provided by Bernoulli and Poisson GCWMs from (3.1) giving parameter pairs (ψ_k, λ_k) where $k \in \{1 \dots K\}$. The second EM procedure then optimizes (3.2). The ZI-GCWM is then compared against the standard Poisson GCWM using a liklihood ratio test which is commented in section 3.6.

3.1 The EM Algorithm for Parameter Estimation

In most finite mixture problems, the standard method for estimating parameters of mixture models is based on the EM algorithm and further discussed by McLachlan and Peel (2000). The Bernoulli-Poisson partitioning method is split into two EM algorithm steps. The first EM partitions the sample space, while the second EM optimizes the zero inflated portion.

3.2 EM - Partitioning

The EM algorithm is based on the local maximum likelihood estimation. The initial values of the parameter estimates can be generated from a variety of strategies outlined in Biernacki et al. (2000), then the algorithm

proceeds by alternation of the E-step and M-step to update parameter estimates. To find an optimal number of components, maximum likelihood estimation is obtained over a range of G, and the best model is selected based on the Bayesian information criterion (BIC).

The convergence criterion of the EM algorithm is based on the Aitken acceleration. It is used to estimate the asymptotic maximum of the log-likelihood at each iteration of the EM algorithm. when the relative increase in the log-likelihood function is no bigger than a small pre-specified tolerance value or the number of iterations reach a limit. In this subsection, we explain the parameter estimation in line with the GCWM methodology proposed by Ingrassia and Minotti (2015). The proposed GCWM a model is based on the assumption that $q(y|\boldsymbol{x}, \vartheta_j)$ belongs to the exponential family of distributions that are strictly related to GLMs. The link function relates the expected value $g(\mu_j) = \beta_{0j} + \beta_{1j}x_1 + \cdots + \beta_{pj}x_p$. We are interested in estimation of the vector $\boldsymbol{\beta}_j$, thus the distribution of $y|\boldsymbol{x}$ is denoted by $q(y|\boldsymbol{x}; \boldsymbol{\beta_j}, \lambda_j)$, where λ_j denotes an additional parameter to account for when a distribution belong to a two-parameter exponential family.

The marginal distribution $p(\boldsymbol{x}; \boldsymbol{\theta}_j)$ has the following components: $p(\boldsymbol{t}; \boldsymbol{\theta}_j^\star)$, $p(\boldsymbol{w}; \boldsymbol{\theta}_j^{\star\star})$, and $p(\boldsymbol{u}; \boldsymbol{\theta}_j^{\star\star\star})$. The first marginal density $p(\boldsymbol{t}; \boldsymbol{\theta}_j^\star)$ is modeled as a Gaussian with mean $\boldsymbol{\mu_j}$ and covariance matrix $\boldsymbol{\Sigma}_j$ as $p(\boldsymbol{t}; \boldsymbol{\mu_j}, \boldsymbol{\Sigma}_j)$. The marginal density $p(\boldsymbol{w}; \boldsymbol{\theta}_j^{\star\star})$ assume that each finite discrete covariate W is represented as a vector $\boldsymbol{w}^r = (w^{r1}, \ldots, \boldsymbol{w}^{rc_r})'$ where $w^{rs} = 1$ if $w_r = s$, such that $s \in \{1, \ldots, c_r\}$, and $w^{rs} = 0$ otherwise.

$$p(\boldsymbol{w}, \boldsymbol{\gamma_j}) = \prod_{r=1}^{d} \prod_{s=1}^{c_r} (\gamma_{jrs})^{w^{rs}} \tag{3.3}$$

for $j = 1, \ldots, G$, where $\boldsymbol{\gamma_j} = (\gamma_{j1}', \ldots, \gamma_{jd}')'$, $\boldsymbol{\gamma}_{jr} = (\gamma_{jr1}', \ldots, \gamma_{jrc_d}')'$, $\gamma_{jrs} > 0$, and $\sum_{s=1}^{c_r} \gamma_{jrs}$, $r = 1, \ldots, q$. The density $p(\boldsymbol{w}, \boldsymbol{\gamma_j})$ represents the product of d conditionally independent multinomial distributions with parameters $\boldsymbol{\gamma}_{jr}$, $r = 1, \ldots, d$. The last marginal density $p(\boldsymbol{u}; \boldsymbol{\theta}_j^{\star\star\star})$ will be modelled as lognormal with mean vector $\boldsymbol{\mu}_j^{\star\star\star}$, and covariance matrix $\boldsymbol{\Sigma}_j^{\star\star\star}$.

Let $(\boldsymbol{x}_1, y_1), \ldots, (\boldsymbol{x}_n, y_n)$ be a sample of n independent observations drawn from model in (2.3). For this sample, the complete data likelihood function, $L_c(\boldsymbol{\Phi})$, is given by

$$L_c(\boldsymbol{\Phi}) = \prod_{i=1}^{n} \prod_{j=1}^{G} \left[\tau_j q(y_i|x_i, \boldsymbol{\beta}_j, \lambda_j) p(t_i, \boldsymbol{\mu}_j, \boldsymbol{\Sigma}_j) p(w_i, \gamma_j) p(u_i, \boldsymbol{\mu_j}^{\star\star\star}, \boldsymbol{\Sigma_j}^{\star\star\star}) \right]^{z_{ij}}, \tag{3.4}$$

where z_{ig} is the latent indicator variable with value of $z_{ig} = 1$ indicating that observation $(\boldsymbol{x_i}, y_i)$, originated from the jth mixture component and $z_{ij} = 0$ otherwise.

By taking the logarithm of (3.4), the complete-data log-likelihood function $\ell_c(\mathbf{\Phi})$ is written by

$$\ell_c(\mathbf{\Phi}) = \sum_{i=1}^{n}\sum_{j=1}^{G} z_{ij}\big[\log(\tau_j) + \log q(y_i|x_i,\boldsymbol{\beta}_j,\lambda_j) + \log p(t_i,\boldsymbol{\mu}_j,\mathbf{\Sigma}_j) + \log p(w_i,\gamma_j)$$

$$+ \log p(u_i,\boldsymbol{\mu}_j^{\star\star\star},\mathbf{\Sigma}_j^{\star\star\star})\big].$$

3.2.1 E-Step - Partitioning

The E-step does not depend on the form of density, and the latent data only relate to $\boldsymbol{z}$. The posterior probability that $(\boldsymbol{x_i}, y_i)$ comes from the jth mixture component is calculated at the sth iteration of the EM algorithm as

$$\pi_{ij}{}^{(s)} = E[z_{ij}|(\boldsymbol{x_i},y_i),\mathbf{\Phi}^{(s)}]$$

$$= \frac{\tau_j{}^{(s)}q(y_i|x_i,\boldsymbol{\beta}_j^{(s)},\lambda_j^{(s)})p(t_i,\boldsymbol{\mu}_j^{(s)},\mathbf{\Sigma}_j^{(s)})p(w_i,\gamma_j^{(s)})p(u_i,\boldsymbol{\mu}_j^{\star\star\star(s)},\mathbf{\Sigma}_j^{\star\star\star(s)})}{f(\boldsymbol{x_i},y_i;\Phi^{(s)})}.$$

3.2.2 M-Step - Partitioning

It follows that at the $(s+1)$th iteration, the conditional expectation of (3.2) on the observed data and the estimates from the sth iteration results in

$$Q(\mathbf{\Phi}|\mathbf{\Phi}^{(s)}) = \sum_{i=1}^{n}\sum_{j=1}^{G}\pi_{ij}^{(s)}\big[\log(\tau_j)+\log q(y_i|x_i,\boldsymbol{\beta}_j,\lambda_j)+\log p(t_i,\boldsymbol{\mu}_j,\mathbf{\Sigma}_j)+\log p(w_i,\gamma_j)+\log p(u_i,\boldsymbol{\mu}_j^{\star\star\star(s)},\mathbf{\Sigma}_j^{\star\star\star(s)})\big]$$

$$= \sum_{i=1}^{n}\sum_{j=1}^{G}\pi_{ij}^{(s)}\log(\tau_j) + \sum_{i=1}^{n}\sum_{j=1}^{G}\pi_{ij}^{(s)}\log q(y_i|x_i,\boldsymbol{\beta}_j^{(s)},\lambda_j^{(s)}) + \sum_{i=1}^{n}\sum_{j=1}^{G}\pi_{ij}^{(s)}\log p(t_i,\boldsymbol{\mu}_j^{(s)},\mathbf{\Sigma}_j^{(s)})$$

$$+ \sum_{i=1}^{n}\sum_{j=1}^{G}\pi_{ij}^{(s)}\log p(w_i,\gamma_j^{(s)}) + \sum_{i=1}^{n}\sum_{j=1}^{G}\pi_{ij}^{(s)}\log p(u_i,\boldsymbol{\mu}_j^{\star\star\star(s)},\mathbf{\Sigma}_j^{\star\star\star(s)}). \quad (3.5)$$

The M-step requires maximization of the Q-function with respect to $\mathbf{\Phi}$ which can be done separately for each term on the right hand side in (3.5). As a result, the parameter updates $\hat{\tau}_j$, $\hat{\boldsymbol{\mu}}_j$, $\hat{\boldsymbol{\sigma}}_j$, and $\hat{\boldsymbol{\gamma}}_j$ on the $(s+1)$th iteration are:

$$\hat{\tau}_j^{(s+1)} = \frac{1}{n}\sum_{i=1}^{n}\pi_{ij}^{(s)}, \qquad \hat{\boldsymbol{\mu}}_j^{(s+1)} = \frac{1}{\sum_{i=1}^{n}\pi_{ij}^{(s)}}\sum_{i=1}^{n}\pi_{ij}^{(s)}\boldsymbol{t}_i, \qquad \hat{\gamma}_{jr}^{(s+1)} = \frac{\sum_{i=1}^{n}\pi_{ij}^{(s)}\omega_i^{rs}}{\sum_{i=1}^{n}\pi_{ij}^{(s)}},$$

$$\hat{\boldsymbol{\sigma}}_j^{(s+1)} = \frac{1}{\sum_{i=1}^{n}\pi_{ij}^{(s)}}\sum_{i=1}^{n}\pi_{ij}^{(s)}(\boldsymbol{t}_i - \hat{\boldsymbol{\mu}}_j^{(s+1)})(\boldsymbol{t}_i - \hat{\boldsymbol{\mu}}_j^{(s+1)})',$$

The log-normal distribution is relevant for modelling actuarial data. Parameter estimates for the log-normal distribution follow similar suit.

$$\hat{\tau}_j^{(s+1)} = \frac{1}{n}\sum_{i=1}^{n}\pi_{ij}^{(s)}, \qquad \hat{\boldsymbol{\mu}}_j^{\star\star\star(s+1)} = \frac{1}{\sum_{i=1}^{n}\pi_{ij}^{(s)}}\sum_{i=1}^{n}\pi_{ij}^{(s)}\ln\boldsymbol{u}_i,$$

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$$\hat{\boldsymbol{\sigma}}_j^{\star\star\star(s+1)} = \frac{1}{\sum_{i=1}^n \pi_{ij}^{(s)}} \sum_{i=1}^n \pi_{ij}^{(s)} (\ln \boldsymbol{u}_i - \hat{\boldsymbol{\mu}}_j^{\star\star\star(s+1)})(\ln \boldsymbol{u}_i - \hat{\boldsymbol{\mu}}_j^{\star\star\star(s+1)})'.$$

The estimates of $\boldsymbol{\beta}$ are computed by maximizing each of the G terms

$$\sum_{i=1}^n \pi_{ij}^{(s)} \log q(y_i | \boldsymbol{x}_i, \boldsymbol{\beta}_j, \lambda_j). \tag{3.6}$$

Maximization of (3.6) is performed by numerical optimization in R software in a similar framework the mixture of generalized linear models are implemented. For additional details about this implementation the reader is refer to Wedel and De Sabro (1995) and Wedel (2002). For insurance applications, current TCWM model can be used for modeling frequency of claims assuming that $\boldsymbol{Y}$ belongs to Poisson or Bernoulli distributions. When modelling severity of claims, $\boldsymbol{Y}$ can be assumed accommodate Gamma or Lognormal distributions. All of these applications are based on CWM as the underlying approach. For additional information, the reader is referred to the manual of the `flexCWM` package manual for **R** users written by Ingrassia and Minotti (2015).

3.3   EM - Zero-inflated

The optimization of the zero-inflated model (3.2) uses the EM algorithm for maximizing the incomplete-data log-liklihood iteratively (Lambert, 1992). The log-liklihood for ψ_k and λ_k is expressed as

$$l(\psi_k, \lambda_k; y, \boldsymbol{x}) = \sum_{y_i=0} \log \left[e^{\tilde{\boldsymbol{x}}_i \bar{\boldsymbol{\beta}}'_{\boldsymbol{k}}} + \exp\left(-e^{-\tilde{\boldsymbol{x}}_i \boldsymbol{\beta}'_{\boldsymbol{k}}}\right) \right] + \sum_{y_i>0} \left(y_i \tilde{\boldsymbol{x}}_i \boldsymbol{\beta}'_{\boldsymbol{k}} + e^{\tilde{\boldsymbol{x}}_i \boldsymbol{\beta}'_{\boldsymbol{k}}} \right) - \sum_{i=1}^n \log \left(1 + e^{\tilde{\boldsymbol{x}}_i \bar{\beta}_k'} \right) - \sum_{y_i>0} \log(y_i!)$$

Where y_i, and $\tilde{\boldsymbol{x}}_i$ referes to the ith row of the response variable y and covariate matrix $\tilde{\boldsymbol{x}}$. Due to the first term, the log-liklihood is complicated to maximize, Lambert (1992) provides a meaningful solution. Suppose that we could observe $Z_{ik} = 1$ when y_i is generated from the Bernoulli random variable of partition k, and $Z_{ik} = 0$ when y_i is generated from the Poisson random variable. Then the complete-data log-liklihood would be written as

$$l_c(\psi_k, \lambda_k; y, \boldsymbol{z_k}) = \sum_{i=1}^n \left(z_{ik} \tilde{\boldsymbol{x}_i} \bar{\boldsymbol{\beta}_k}' - \log(1 + e^{\tilde{\boldsymbol{x}}_i \bar{\boldsymbol{\beta}_k}'}) \right) + \sum_{i=1}^n (1 - z_{ik})(y_i \tilde{\boldsymbol{x}_i} \boldsymbol{\beta}'_{\boldsymbol{k}} - e^{\tilde{\boldsymbol{x}_i} \boldsymbol{\beta}'_{\boldsymbol{k}}}) + \sum_{i=1}^n (1 - z_{ik}) \log(y_i!)$$

$$= \quad l_c(\psi_k; \boldsymbol{y}, \boldsymbol{z_k}) + l_c(\lambda_k; \boldsymbol{y}, \boldsymbol{z_k}) + \sum_{i=1}^n (1 - z_{ik}) \log(y_i!) \tag{3.7}$$

where z_{ik} is a realization of Z_{ik}. (3.7) is easier to maximize since $l_c(\psi_k; \boldsymbol{y}, \boldsymbol{z_k})$ and $l_c(\lambda_k; \boldsymbol{y}, \boldsymbol{z_k})$ can be maximized seperately for parameters λ_k and ψ_k. With the EM algorithm, the incomplete-data log-liklihood can be maximized iteratively between estimating Z_{ik} with its expectation under current parameters λ_k and ψ_k

(E-Step) and then maximizing the complete data-logliklihood (M-Step).

3.3.1 E-step - Zero-inflated

Using current estimates $\psi_k^{(s)}$ and $\lambda_k^{(s)}$ from the partition Ω_{Z_k}, we calculate the expected value of Z_{ik} by its posterior mean $z_{ik}^{(s)}$ for each cluster k, at iteration s

$$z_{ik}^{(s)} = \begin{cases} \left[1 + \exp\left(-\tilde{\boldsymbol{x}_i}\bar{\boldsymbol{\beta}_k}^{'(s)} - e^{\tilde{\boldsymbol{x}_i}\beta_k^{'(s)}}\right)\right]^{-1}, & y_i = 0 \\ 0, & y_i > 0. \end{cases}$$

3.3.2 M-Step - Zero-inflated

The M-Step can be split into the maximization of two complete data log-likelihoods and the $\boldsymbol{z_i}$ calculated from the previous iteration (s) as:

$$l_c(\psi_k; \boldsymbol{y}, \boldsymbol{z_k^{(s)}}) = \sum_{i=1}^{n} \left(z_{ik}^{(s)}\tilde{\boldsymbol{x}_i}\bar{\boldsymbol{\beta}_k}^{'(s)} - \log(1 + e^{\tilde{\boldsymbol{x}_i}\bar{\boldsymbol{\beta}_k}^{'(s)}})\right) \tag{3.8}$$

$$l_c(\lambda_k; \boldsymbol{y}, \boldsymbol{z_k^{(s)}}) = \sum_{i=1}^{n} (1 - z_{ik}^{(s)})(y_i\tilde{\boldsymbol{x}_i}\boldsymbol{\beta}_k^{'(s)} - e^{\tilde{\boldsymbol{x}_i}\beta_k^{'(s)}}) \tag{3.9}$$

The maximization of (3.9) for GLM coefficients λ_k can be found by using a weighted, log-linear Poisson regression with weights $1 - z_{ik}^{(s)}$ (McCullagh and Nelder, 1989), yielding $\lambda_k^{(s+1)}$. While the parameter for (3.9) can be maximized over a gradient (Lambert, 1992) yielding $\psi_k^{(s+1)}$.

3.4 Comparing zero-inflated Models

Until recently the usual test for comparing zero-inflated to non-zero inflated models has been the Vuong Test for non-nested models (see Vuong (1989)). However recent work has shown the misuse of this test for zero inflation (see Wilson (2015)). The Wilson and Einbeck (2018) show that it is sufficient to test for zero-modificiation in the form of a liklihood ratio test. The test is defined as follows

$$H_0: \quad \psi_k = 0 \qquad vs. \qquad H_a: \quad \psi_k \neq 0$$

$$\varphi = -2\left[l(\tilde{\lambda}_k; y, \boldsymbol{x}) - l(\lambda_k, \psi_k; y, \boldsymbol{x})\right]$$

where $l(\tilde{\lambda}_k; y, \boldsymbol{x})$ is the log-liklihood of a single component GCWM Poisson model on Ω_{Z_k} parameterized by $\tilde{\lambda}_k$. Recall that ψ_k is the zero-inflation paramater of the kth parition. The test statistic φ is shown to be distributed Chi-square with m degrees of freedom (χ_m^2) and $\alpha = 0.10$ (Wilson and Einbeck, 2018).

4 Application

4.1 Data

We illustrate the proposed methodology on French motor claims data set by policy. This data set is available as part of the R package `CASdatasets` and it is previously used in the book *Computational Actuarial Science with R* by Charpentier (2014). The book demonstrated various GLM modeling approaches for fitting frequency and severity of this data. The claim count including zero claims for 413,169 motor third-party liability policies are provided with the associated risk characteristics. The loss amounts by policy ID are also provided.

Table 1: The description of variables in the French Motor Third-Part Liability dataset.

| Attribute	Description
Policy ID	Unique identifier of the policy holder
Claim Nb	Number of claims during exposure period (0,1,2,3,4)
Exposure	The exposure of policy in years (0–1.5)
Power	Power level of car ordered categorical (12 levels)
Car Age	Car age in years
Driver Age	Age of a legal driver
Brand	Car brands (7 types)
Gas	Diesel or Regular
Region	Regions in France (10 classifications)
Density	Number of inhabitants per km^2
Loss Amount	Portion of claim the insurance policy pays

4.2 Analysis and Results

4.2.1 Modelling Severity

In this section, we show the results from modeling French motor losses. We consider the following covariates: density, driver age, car age, power, gas, and region. The model that was fitted is defined with the following equation where $\epsilon \sim \mathcal{N}(0, \sigma)$, and

$$LossAmount = Density + CarAge + DriverAge + Region + Power + Gas + \epsilon. \qquad (4.1)$$

The canonnical log-link is used for the GLM (4.1). Similarly to Miljkovic and Fernndez (2018), car age is modelled as a categorical variable with five categories: $[0, 1), [1, 5), [5, 10), [10, 15),$ and 15+. Additionaly, driver age is modelled as a categorical variable with five categories: $[18, 23), [23, 27), [27, 43), [43, 75),$ and 75+. Power is modelled into three categories as in Charpentier (2014): DEF, GH, and other.

Figure 1: Density variable: Left figure shows the fit when Gaussian distribution is imposed (CMW approach) to highly skewed data. Right figure shows the fit when log-normal transformation is applied (GL-TCWM approach).
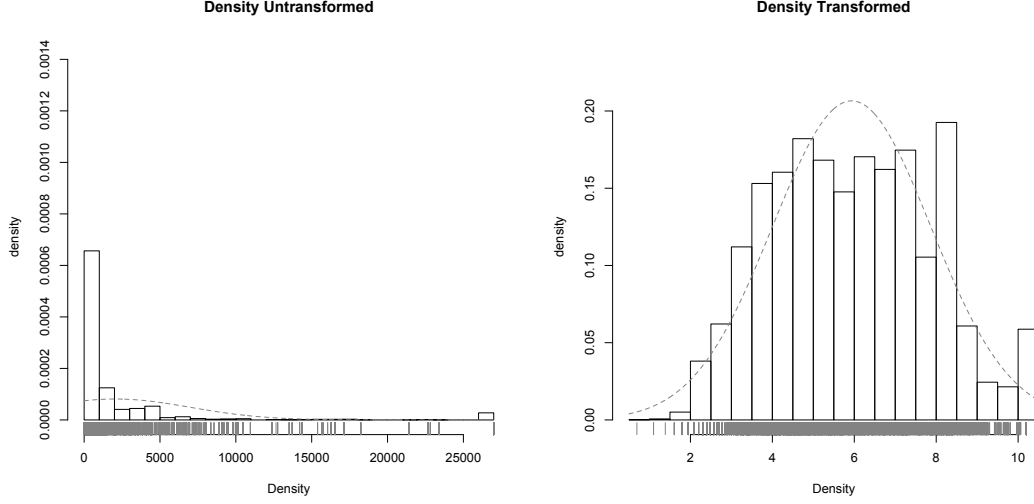
Beginning with the continuous covariate $Density$, we want to inspect the shape of its univariate data to see if it follows Gaussian distribution. The left side of Figure 2 clearly revels that the the $Density$ is rather skewed right with several observations that report high value of density. This indicates a need for a transformation. With the log-normal assumption, the $Density$ is transformed which improves the fit (see the right side of Figure 2) on the data.

Table 2: Comparing GCWM vs CWM models.

Model	k	AIC	BIC
CWM	1	352470	352661
	2	314560	314949
	3	301223	301812
	4	287020	287808
	5	**284283**	**285268**
GCWM	1	111129	111320
	2	90039	90428
	3	89476	90065
	4	**88781**	**89568**
	5	88731	89717

The result of the transformation is a better AIC and BIC. Table 2 shows a considerable difference in BIC

and AIC comparing CWM and GCWM. The five component CWM with a BIC of 285268 is significantly higher than the four component GCWM with a considerably lower BIC of 89568.

4.2.2 Driver Analysis of Severity Model

We now investigate the results of GCWM in relation to the valuation of risk. For practical uses, finding clusters allows us to create different classifications of risk for various fields of drivers. The following GCWM allows to cluster different drivers in groups allowing one to assign different rates to different clusters.
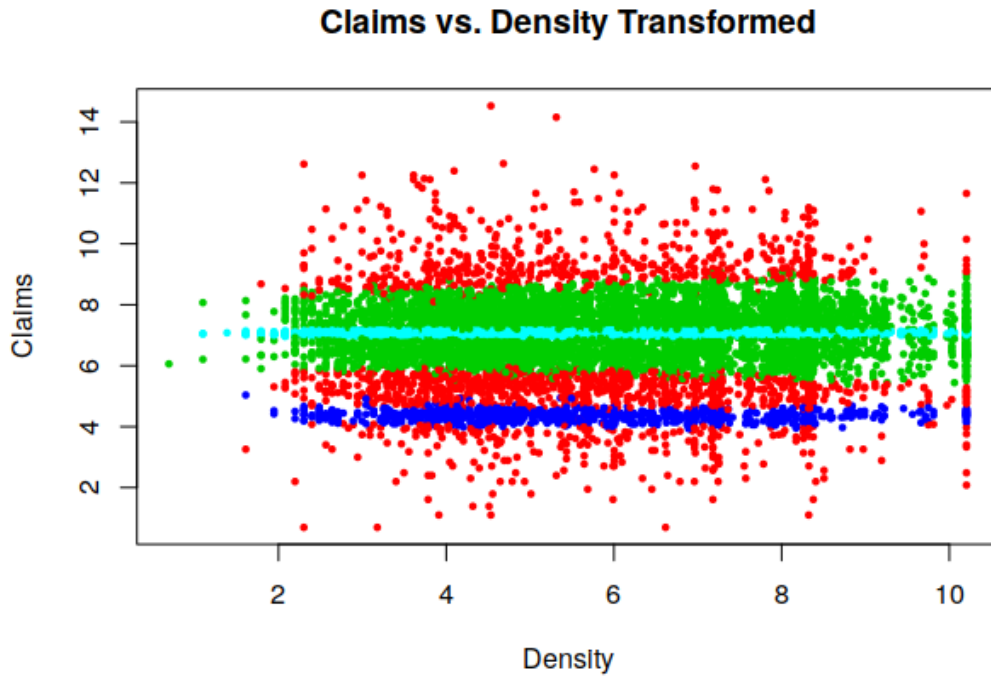

Figure 2: Showing clusters in color for Claims vs Density under lognormal assumptions.

Table 3: Size of clusters for the GCWM a model.

1	2	3	4
1683	5766	848	7093
Red	Green	Blue	Teal

After fitting the model, we then take a look at the size of each cluster. The GCWM a function has chosen four components as the best model to represent the data. The size of each cluster is displayed in Table 3. Attention is brought to largest quantity of drivers that are grouped into cluster 4. This accounts for 46% of all

drivers and is fairly concentrated in the center of Figure 2. From the results we can create an insurance model with the following characteristics. Cluster 3 drivers have both low variability and low average cost in claims, thus can be insured with a lower rate than other drivers. From a risk management perspective this is the most ideal case as these claims have very low variance and cost. Cluster 4 drivers have also low variability but a higher average cost, thus they would have a rate higher than cluster 3. Cluster 2 drivers have the next highest cost and variability of the clusters, these drivers are colored in green in Figure 2. The final cluster colored in red has the highest cost and variaiblity in claims out of any of the other clusters. From a risk management perspective this cluster would have the highest rate.

Table 4: Summarized volatility information of each cluster for Claims.

Volatility Level - (Cluster)	Minimum	Mean	Maximum	$\sigma(0.05)$
V1 - (3)	51	79	154	**13.19**
V2 - (4)	1039	3109	1324	**52.07**
V3 - (2)	221	1687	8841	**1284.41**
V4 - (1)	2	9717	2036833	**64835.96**

Table 4 shows a breakdown of the types of drivers, ordered by volatility in descending order. Beginning with V1, these drivers tend to have claims between \$51 to \$154, with a standard deviation of \$13.19, and a mean of \$79. That means that these drivers rarely exceed costs and tend to have very low volatility. Moving onto V2, these drivers have the second level of volatility. Drivers in this range tend to have claims anywhere between \$1039 to \$3109, with a standard deviation of \$ 52.07, and a mean of \$3109. Proceeding to V3, its volatility in claims is greater than the preceding levels. Drivers in this cluster have claims anywhere between \$221 to \$8841, with a mean of \$1687, and a standard deviation of \$1284.41. Finally V4 denotes the level of highest volatility. Claims in this level reach the highest recorded claim of \$ 2036833, a mean of \$9717, and a standard deviation of \$64835.96.

Coefficients of clustered results are used to calculate premiums in car insurance. Table 5 shows the coefficients of the fitted model. The significance codes are defined as ≈ 0 (***), 0.001 (**), 0.01 (*), 0.05 (.) pertaining to the p value of the specific coefficient. In each cluster significance varies but overall the majority of coefficients are significant. An interpretation of the results is as follows. The first cluster (V1) has significant coefficients of Region (R#), Power (P#), Car age (C#), and Driver age. We see that the sign of coefficients change depending on the relative center of each cluster. The comparison between driver age in V1 and V4 is a perfect example. The coefficient of driver age is positive in V1 which shows that with larger age, claims tend to go up. Comparing that with last cluster, (V4). The driver age coefficient is negative. This is due to the fact that ages in V4 are very young (mean age of 25). Thus as age increases drivers tend to take less risks and

Table 5: The coefficients and significance of each cluster.

Coef	V1 (Red)			V2 (Blue)			V3 (Teal)			V4 (Green)		
	Estimate	Error	P	Estimate	Error	P	Estimate	Error	P	Estimate	Error	P
(Intercept)	$7.07 \cdot 10^0$	1.00	***	$6.96 \cdot 10^0$	1.05	***	$5.92 \cdot 10^0$	1.12	***	$8.91 \cdot 10^0$	1.13	***
Density	$1.14 \cdot 10^{-4}$	1.00		$-7.03 \cdot 10^{-3}$	1.00	*	$-3.47 \cdot 10^{-2}$	1.01	***	$3.03 \cdot 10^{-2}$	1.01	***
DriverAge	$1.03 \cdot 10^{-4}$	1.00	***	$9.89 \cdot 10^{-4}$	1.00	.	$1.88 \cdot 10^{-2}$	1.00	***	$-6.39 \cdot 10^{-2}$	1.00	***
R23	$4.94 \cdot 10^{-3}$	1.00		$-1.39 \cdot 10^{-1}$	1.04	***	$1.47 \cdot 10^{-1}$	1.12		$-1.91 \cdot 10^{-1}$	1.11	.
R24	$-1.43 \cdot 10^{-2}$	1.00	***	$6.58 \cdot 10^{-2}$	1.02	***	$-3.61 \cdot 10^{-1}$	1.05	***	$-1.71 \cdot 10^{-2}$	1.04	
R25	$-1.43 \cdot 10^{-2}$	1.00	***	$-1.78 \cdot 10^{-1}$	1.04	***	$-1.05 \cdot 10^{-1}$	1.09		$1.59 \cdot 10^{-1}$	1.09	.
R31	$-2.32 \cdot 10^{-3}$	1.00		$2.43 \cdot 10^{-3}$	1.03		$1.98 \cdot 10^{-1}$	1.07	**	$-8.98 \cdot 10^{-2}$	1.06	.
R52	$-1.52 \cdot 10^{-2}$	1.00	***	$2.23 \cdot 10^{-1}$	1.02		$-3.89 \cdot 10^{-1}$	1.06	***	$1.36 \cdot 10^{-1}$	1.06	*
R53	$-1.38 \cdot 10^{-2}$	1.00	***	$2.22 \cdot 10^{-1}$	1.02	***	$-1.78 \cdot 10^{-1}$	1.06	**	$8.09 \cdot 10^{-2}$	1.06	
R54	$-1.47 \cdot 10^{-2}$	1.00	***	$8.69 \cdot 10^{-4}$	1.03	***	$-4.04 \cdot 10^{-1}$	1.08	***	$1.85 \cdot 10^{-1}$	1.07	**
R72	$-5.98 \cdot 10^{-3}$	1.00	***	$1.44 \cdot 10^{-1}$	1.03	***	$-6.77 \cdot 10^{-2}$	1.07		$2.37 \cdot 10^{-2}$	1.06	
R74	$-1.82 \cdot 10^{-2}$	1.00	***	$1.44 \cdot 10^{-1}$	1.06		$-1.05 \cdot 10^{-1}$	1.14		$-3.32 \cdot 10^{-1}$	1.14	*
PE	$-1.21 \cdot 10^{-3}$	1.00		$3.31 \cdot 10^{-2}$	1.02	***	$6.47 \cdot 10^{-2}$	1.05		$-1.83 \cdot 10^{-1}$	1.04	***
PF	$-2.56 \cdot 10^{-3}$	1.00	*	$7.85 \cdot 10^{-2}$	1.02	.	$1.43 \cdot 10^{-1}$	1.04	**	$-3.93 \cdot 10^{-2}$	1.04	
PG	$-5.31 \cdot 10^{-4}$	1.00		$1.83 \cdot 10^{-1}$	1.02	***	$1.35 \cdot 10^{-1}$	1.05	**	$-7.17 \cdot 10^{-2}$	1.04	.
PH	$3.89 \cdot 10^{-3}$	1.00	*	$9.02 \cdot 10^{-2}$	1.03	***	$1.15 \cdot 10^{-1}$	1.07	.	$-1.83 \cdot 10^{-1}$	1.07	**
PI	$-2.00 \cdot 10^{-3}$	1.00		$1.20 \cdot 10^{-1}$	1.03	**	$2.04 \cdot 10^{-1}$	1.08	**	$-8.05 \cdot 10^{-2}$	1.07	
PJ	$8.36 \cdot 10^{-3}$	1.00	***	$-5.91 \cdot 10^{-2}$	1.03	***	$2.54 \cdot 10^{-1}$	1.08	***	$-1.61 \cdot 10^{-1}$	1.08	*
PK	$5.63 \cdot 10^{-3}$	1.00	*	$6.86 \cdot 10^{-2}$	1.04		$1.47 \cdot 10^{-1}$	1.09	.	$-1.80 \cdot 10^{-1}$	1.12	
PL	$9.04 \cdot 10^{-3}$	1.00	**	$4.18 \cdot 10^{-1}$	1.06		$2.31 \cdot 10^{-1}$	1.14	.	$3.09 \cdot 10^{-1}$	1.18	.
PM	$-7.52 \cdot 10^{-3}$	1.01		$1.14 \cdot 10^{-1}$	1.08	***	$-7.67 \cdot 10^{-2}$	1.20		$1.68 \cdot 10^{-1}$	1.28	
PN	$-3.92 \cdot 10^{-3}$	1.01		$-5.77 \cdot 10^{-1}$	1.09		$-4.75 \cdot 10^{-1}$	1.24	*	$-1.58 \cdot 10^{-1}$	1.21	
PO	$3.48 \cdot 10^{-3}$	1.01		$3.57 \cdot 10^{-2}$	1.09	***	$1.01 \cdot 10^{-1}$	1.23		$-6.48 \cdot 10^0$	1.63	***
C1	$5.67 \cdot 10^{-3}$	1.00	*	$1.83 \cdot 10^{-1}$	1.03		$3.74 \cdot 10^{-2}$	1.08		$-2.34 \cdot 10^{-1}$	1.08	**
C2	$4.58 \cdot 10^{-3}$	1.00	*	$9.89 \cdot 10^{-4}$	1.03	***	$-1.02 \cdot 10^{-1}$	1.07		$-5.90 \cdot 10^{-1}$	1.07	***
C3	$8.08 \cdot 10^{-3}$	1.00	***	$1.44 \cdot 10^{-1}$	1.03	***	$-1.98 \cdot 10^{-1}$	1.07	**	$-5.99 \cdot 10^{-1}$	1.07	***
C4	$8.25 \cdot 10^{-3}$	1.00	***	$2.76 \cdot 10^{-1}$	1.03	***	$-2.21 \cdot 10^{-1}$	1.08	**	$-3.87 \cdot 10^{-1}$	1.09	***
C5	$4.09 \cdot 10^{-3}$	1.00	.	$2.09 \cdot 10^{-1}$	1.03	***	$-2.13 \cdot 10^{-1}$	1.09	**	$-9.27 \cdot 10^{-1}$	1.09	***

become more responsible, which would decrease cost of claims. We see the same interpretation with Density. Clusters V2, V3, and V4 have density as a significant coefficient. However, the sign of the coefficient changes to negative in V4, while V2 and V3 have positive coefficients. Again this due to the relative centres of the cluster indicate both the magnitude and sign of the coefficient.

To summarize, the drivers have been clustered into four categories with distinct characteristics outlined in Table 4. We have seen how using the results from GCWM a, one can create an insurance model based on clustering algorithms with various levels of risk represented in each cluster. An interesting result was that cluster 1 contradicted the usual understanding that young people take more risks. The GCWM a method found a group that was the clear majority of drivers, in which the volatility of their claims was extremely low regardless of Driver Age, or Density. This finding shows that GCWM a may potentially find unique models that are

otherwise hidden within the data.

4.2.3 Modeling Claims Frequency

In this section, we model frequency of the French motor claims. We consider the covariates log-density, driver age, car age, and a three-class grouping of the power of a car labelled Power F. The choice of covariates stems from the previously modelled single component ZIP (Charpentier, 2014). For the purposes of computational feasibility, only claims from the largest populated insurance region (R24) had been selected. The extension into multiple components is modelled with the linear formula

$$\text{Claim Nb} \sim \text{Driver Age} + \text{Log Density} + \text{Car Age} + \text{Power F}.$$

After fitting the model, the size of each cluster is noted. The GCWM a has chosen 3 components as the best model to represent the data. The size of each cluster is displayed in Table **??**. Attention is brought to cluster 2 which holds nearly 67% of the total population. Cluster 3 holds the fewest amount of drivers with merely 2.4% of the total population. Table 4 shows an in-depth analysis of Driver Age and Claim Number as shown in 2. Cluster 1 has the youngest drivers of the whole population with a mean age of 29.39 and a standard deviation of 4.87. Cluster 2 has a relatively older age group with a mean age of 36.90, and a standard deviation of 1.20. Finally Cluster 3 shows the oldest age group of drivers with a mean age of 54.25 and a standard deviation of 11.66. However when looking at the relative proportion of claims for each cluster as shown in Table 6. One notices that the middle age group has a higher proportion of non-zero claims. Relative to the other clusters, the middle age group has a non-zero claim proportion of 17.25 %. This is bolded in the Table 6 to show the difference of cluster 2 in comparison to 1, and 3.
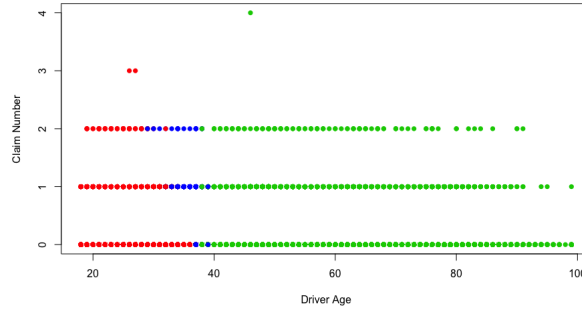

Figure 3: Claim number versus driver age, with partitions denoted by colour.

The significance of the codes are defined as ≈ 0 (***), 0.001 (**), 0.01 (*), 0.05 (.) pertaining to the p value of the specific coefficient. For each cluster the significance of the coefficients vary greatly. In cluster

Table 6: Claim Nb analysis of clusters

Cluster	Claim Nb	Counts	Proportion (%)
1	0	48172	96.98 %
	1	1432	2.90 %
	2	64	0.12 %
	3	2	$\approx 0\%$
2	0	3262	**82.75 %**
	1	658	**16.69 %**
	2	22	**0.56 %**
3	0	102905	96.18%
	1	3963	3.71 %
	2	120	0.11 %
	4	1	$\approx 0\%$

Table 7: Size of clusters for modelling claims

1	2	3
49670	106989	3942
Red	Green	Blue

Table 8: Driver Age analysis of clusters.

Cluster	Minimum	Maximum	Mean	σ
1	18	38	29.39	4.87
2	28	39	36.90	1.20
3	38	99	54.25	11.66

1, the count parameters (β_{xn}) for the Intercept, Driver Age and Log-Density are highly significant (***). The zero parameters β_{xz} are all highly significant with the exception of the power group GH (β_{5z}). After comparing the zero-inflated model for cluster 2 with a reduced Poisson model, the Vuong statistic had chosen the simpler Poisson model shown in Table 9. The Poisson model shows significance for the Intercept, Car Age, and Log-Density. Cluster 3 shows signficance for both zero and count models. The Intercept, Driver Age, and Log Density for both models show high significance. In summary the drivers have been clustered into three categories outlined in Table 9. From the coefficients, one can generate a premium plan tailored to the specific classifications of drivers using GCWM a.

5   Simulation Study

Two simulation studies are conducted to determine the validity of transformation and the effectiveness of the Bernoulli-Poisson partitioning method. The first section outlines the need for transformation in the covariates. The second section shows the classification accuracy and other relevant analysis for the Bernoulli-Poisson method.

Table 9: Table of coefficients for each cluster

Par	Est.	Std. Err	Z Val.	P	Est.	Std. Err	Z Val.	P	Est.	Std. Err	Z Val.	P
β_{0n}	-1.621	0.213	-7.629	***	-3.581	0.102	-35.109	***	6.973	0.923	7.553	***
β_{1n}	-0.075	0.006	-11.631	***	0.000	0.001	0.012		-0.220	0.028	-7.773	***
β_{2n}	-0.006	0.005	-1.181		-0.015	0.003	-5.481	***	0.003	0.009	0.325	
β_{3n}	0.113	0.015	7.589	***	0.085	0.010	8.884	***	0.103	0.030	3.401	***
β_{4n}	-0.065	0.086	-0.752		0.094	0.054	1.611		-0.087	0.153	-0.570	
β_{5n}	0.081	0.092	0.875		0.074	0.058	1.287		-0.006	0.167	-0.039	
β_{0z}	-228.080	55.429	-4.115	***					-106.767	6.873	-15.534	***
β_{1z}	7.601	1.832	4.149	***					3.059	0.194	15.782	***
β_{2z}	-0.313	0.128	-2.447	*					0.017	0.019	0.921	
β_{3z}	-4.309	1.002	-4.303	***					-1.133	0.087	-12.967	***
β_{4z}	7.891	2.105	3.750	***					-0.094	0.313	-0.301	
β_{5z}	-1.267	1.346	-0.941						0.096	0.334	0.288	

5.1 Simulation Study - Transformation

In this section, we show how the proposed methodology works for different simulation settings. The simulation study was generated based on the regression coefficients of the **CASdataset** used in the previous section. The aim of the simulation study was to test the accuracy and ability of both GCWM a and CWM to return estimates of true parameters when one or more of the covariates is lognormal and the other two are Gaussian. This was designed to test both functions in the event when one of the covariates is non-Gaussian. The motivation behind this is fact is that many covariates used in insurance are likely to come from non-Gaussian distribution. Thus this was aimed to test the relevancy of CWM, which treats all covariates as Gaussian.

We define Model 1 as the base line model in which the coefficients were generated for **CASdataset** and reported in upper portion of Table 2. These coefficients were then rounded and treated as true parameters. A simulation with three GLM mixture components was then generated around these true parameters in which the third covariate X_3 was lognormal. Stemming from this, both CWM and GCWM a were run. The GCWM a treats X_3 as a lognormal covariate which then applies the relevant transformation.

The results for Model 1 were summarized in upper portion of Table 3 based on the performance of the GCWM a approach. The simulation was run 1000 times. We reported the percentage of runs for each predictor and the corresponding intercept in each mixture component under the assumption of 5% error. For example, predictor X_2 in the component 2 of Model 1 reported 90.10% accuracy. This means that 90.1% of the time the true parameter was estimated within 5% error. In this setting, predictor X_1 in the second component was insignificant in the real data set. The purpose of including this parameter in Model 1 was to test the sensitivity of GCWM a for insignificant predictors. In this case, the result of zero is underlined and it means that it has

no influence on the response variable in this simulation. Further, we created Models 2, 3, 4 and 5 by altering the parameters of Model 1 by $+30\%$, -30%, $+50\%$, and -50% accordingly and keeping the second covariate of the second component as an insignificant predictor form the **CASdataset** model. This was done to test the accuracy of GCWM a to the sensitivity of coefficients. Based on the results in Table 3, we can see that GCWM a performs well for all simulation settings.

Table 10: GCWM a vs CWM Accuracy: Covariate X_3 is treated as log-normal, the rest are Gaussian covariates. The transformation of X_3 is considered.

Model	Component	Intercept	X_1	X_2	X_3	Intercept	X_1	X_2	X_3
1	1	93.00%	90.10%	93.00%	93.10%	0.00%	0.00%	0.00%	0.00%
	2	90.10%	<u>0.00%</u>	90.10%	90.10%	0.00%	0.00%	0.00%	0.00%
	3	99.20%	99.10%	99.20%	99.20%	0.00%	0.00%	0.00%	0.00%
2	1	89.80%	89.20%	89.80%	89.80%	0.00%	0.00%	4.60%	0.00%
	2	89.20%	<u>0.00%</u>	89.20%	89.20%	0.00%	<u>0.00%</u>	0.00%	0.00%
	3	99.20%	99.20%	99.20%	99.20%	0.00%	0.20%	1.70%	0.00%
3	1	100.00%	100.00%	100.00%	100.00%	0.00%	0.00%	0.00%	0.00%
	2	100.00%	<u>0.00%</u>	100.00%	100.00%	0.00%	0.00%	0.00%	0.00%
	3	99.20%	99.20%	99.20%	99.20%	0.00%	0.00%	0.00%	0.00%
4	1	88.60%	86.80%	88.60%	87.00%	0.00%	0.00%	0.00%	0.00%
	2	86.90%	<u>0.00%</u>	86.90%	86.90%	0.00%	<u>0.00%</u>	0.00%	0.00%
	3	99.20%	99.20%	99.20%	99.20%	0.00%	0.00%	0.00%	0.00%
5	1	85.90%	84.90%	85.60%	85.90%	0.00%	0.00%	0.00%	0.00%
	2	85.00%	<u>0.00%</u>	84.90%	84.90%	0.00%	<u>0.00%</u>	0.00%	0.00%
	3	99.20%	99.20%	99.20%	99.20%	0.00%	0.20%	10.90%	0.00%

Table 4 provides the summary of the results when CWM was used in the analysis of the same models considered in Table 3. It is not surprising to see that barely any of the simulation runs estimated correctly all parameters as most of the results are zero. This means that the performance of CWM approach is poor in presence of one non-Gaussian covariate which in this case is a log-normal covariate. Similarly to Table 3, Table-4 shows the underlined results pointing to insignificant predictors.

Table 5 provides the summary of Mean Squared Errors (MSE) of each parameter of the models in Table 3 estimated via 1000 simulation runs. The MSE is computed using the following formula $\text{MSE}(\beta_i) = \frac{\sum_i^n (\beta_i - \hat{\beta}_i)^2}{n}$. The MSEs related to the predictor variables for all models and their corresponding components are about zero indicating that GCWM a approach performs well. This is also a result of having a small size coefficients.

Table 6 provides the summary of Mean Squared Errors (MSE) of each parameter of the models in Table 4 estimated via 1000 simulation runs. In contrary to the results reported in Table 5, these results in Table

Table 11: GCWM a results: the summary of MSE for all parameters used in five models. The covariate X_3 is treated as log-normal a nd the rest are Gaussian. These results correspond to those in Table 3.

Model	Component	β_o	MSE(β_o)	β_1	MSE(β_1)	β_2	MSE(β_2)	β_3	MSE(β_3)
1	1	1028	(11.353)	0.03	(0.00)	3.5	(0.00)	-380	(0.09)
	2	1600	(0.000)	-0.01	(0.00)	1.5	(0.00)	-250	(0.00)
	3	40000	(0.035)	-6.00	(0.00)	-305	(0.00)	1100	(0.47)
2	1	1350	(0.167)	0.04	(0.00)	4.5	(0.00)	-500	(0.03)
	2	2080	(0.001)	0.04	(0.00)	2.0	(0.00)	-325	(0.00)
	3	52000	(0.012)	-8.00	(0.00)	450	(0.00)	14300	(0.01)
3	1	720	(0.001)	0.02	(0.00)	2.5	(0.00)	-266	(0.00)
	2	1100	(0.008)	0.00	(0.00)	1.1	(0.00)	-17511	(0.00)
	3	28000	(0.002)	-4.20	(0.00)	245	(0.00)	7700.	(0.00)
4	1	1650	(13.056)	0.05	(0.00)	5.3	(0.00)	-570	(0.00)
	2	2400	(0.000)	-0.01	(0.00)	2.3	(0.00)	-375	(0.00)
	3	60000	(0.051)	-9.00	(0.00)	-457	(0.00)	16500	(0.00)
5	1	500	(1.115)	0.02	(0.00)	2.0	(0.00)	-190	(0.05)
	2	800	(0.003)	0.00	(0.00)	0.8	(0.00)	-120	(0.00)
	3	20000	(0.000)	-3.00	(0.00)	-150	(0.00)	5500	(0.00)

Table 12: CWM results: the summary of MSE for all parameters used in five models. All three covariates are treated as Gaussian. These results correspond to those in Table 4.

Model	Component	β_o	MSE(β_o)	β_1	MSE(β_1)	β_2	MSE(β_2)	β_3	MSE(β_3)
1	1	1028	(·)	0.03	(·)	3.5	(·)	-380	(·)
	2	1600	(·)	-0.01	(·)	1.5	(·)	-250	(·)
	3	40000	(·)	-6.00	(·)	-305	(·)	1100	(·)
2	1	1350	(·)	0.04	(·)	4.5	(·)	-500	(·)
	2	2080	(·)	0.04	(·)	2.0	(·)	-325	(·)
	3	52000	(·)	-8.00	(0.006)	450	(44.1)	14300	(·)
3	1	720	(·)	0.02	(·)	2.5	(·)	-266	(·)
	2	1100	(65.814)	0.00	(·)	1.1	(·)	-17511	(·)
	3	28000	(·)	-4.20	(·)	245	(·)	7700.	(·)
4	1	1650	(·)	0.05	(·)	5.3	(·)	-570	(·)
	2	2400	(·)	-0.01	(·)	2.3	(·)	-375	(·)
	3	60000	(·)	-9.00	(·)	-457	(·)	16500	(·)
5	1	500	(·)	0.02	(·)	2.0	(·)	-190	(·)
	2	800	(·)	0.00	(·)	0.8	(·)	-120	(·)
	3	20000	(·)	-3.00	(0.003)	-150	(4.7)	5500	(·)

6 are significantly different. We can observe that the MSEs for most of the Models and their corresponding components are not generated at all and as such they are shown as (·). This is not surprising because Table 4 shows the accuracy of CWM is not good when attempting to model non-Gaussian predictors as Gaussian.

In summary, our simulation results showed good performance of GCWM a approach in modeling non-Gaussian covariates. More specifically, these results show high accuracy when covariates are log-normal. In contrary, CWM fails to estimate parameters accurately when the Gaussian assumption is violated.

5.2   Simulation Study - Bernoulli-Poisson Partitioning

In this section we show how the Bernoulli-Poisson partitioning (BP) method behaves under different conditions. The components were genereated under similar coefficients taken from the **CASDatasets** package. The coefficients were rounded and treated as true parameters to which data was generated from. The mean and standard deviation of the covariates within each component was also taken into account when generating data. The first simulation examines the performance of the GCWM a model for classification. We generate three components each with sample size $N = 1000$ for a total of 3000 simulated points. The model generated is similar to the mean and standard deviations of Table 8. Consider three simulated covariates and the following GLM model

$$\text{SimClaims} \sim \text{SimDriverAge} + \text{SimLogDensity} + \text{SimCarAge}.$$

Here the GCWM a model is fitted to the simulated data and used to classify into three components. The misclassification rate is calculated by the proportion of true labels placed in other components by the GCWM a model. The results of the simulation is based on the generated dataset are presented in Table 8. The total misclassification rate is 1.8% and the majority of misclassified components are between components two and three.

Table 13: Misclassfication rate and label comparison of generated data.

True Labels	Classified			Misclassification Rate (%)
	1	2	3	
1	992	3	5	0.80
2	0	990	10	1.00
3	15	20	965	3.50
Overall Misclassification Rate				1.80 %
Average Purity				98.23 %
Adjusted Rand Index				0.9479

$$n_{ij} = \text{ across diagonal}, \qquad a_i = \text{ row sums}, \qquad b_j = \text{ column sums}$$

$$ARI = \frac{\sum_{ij} \binom{n_{ij}}{2} - [\sum_i \binom{a_i}{2} \sum_j \binom{b_j}{2}]/\binom{n}{2}}{\frac{1}{2}[\sum_i \binom{a_i}{2} + \sum_j \binom{b_j}{2}] - [\sum_i \binom{a_i}{2} \sum_j \binom{b_j}{2}]/\binom{n}{2}} \qquad AP = \frac{1}{N} \sum_i n_{ij}$$

The experiment is expanded further to show how Bernoulli-Poisson partitioning behaves over 1000 runs and under two different conditions. The first condition is defined as follows. The mean and standard deviations are taken as given by the estimated ZIP components from the **CASDataset**. The second condition involves adjusting the means of two of the covariates so they are closer to each other. The goal is to show that the BP-method holds its use even when means among covariates are close. Conditions are divided into two categories. N is considered normal, where the covariate means are taken directly from the sample data. C is considered to be "close", where the covariate means are manipulated so that they are closer to each other within some degree. This is a common problem in classification where if the means among two different components are close, then misclassification rate increases [Lim et al. (2014)]. Experiment 2 defines the use of 3 different partitioning methods to initialize a zero-inflated model. Poisson method assumes that the presence of non-zeros will provide a better partitioning of the data-set. Bernoulli assumes that the presence of excess zeros will determine the best partitioning of the data-set . Finally the BP-Method assumes that both methods are weighed equally and therefore both must be taken into account when partitioning the dataset. The mean and standard deviation of each measurement is provided in Table 14.

Table 14: Experiment 2: mean and standard deviations for each statistic comparing each method.

Type	Condition	Poisson (σ)	Bernoulli (σ)	BP-Method (σ)
Misclassification Rate	N	1.70% (6.00)	1.60% (6.00)	1.10% (0.02)
	C	5.00% (7.00)	6.00% (2.00)	7.00% (4.00)
Average Purity	N	98.87% (2.00)	98.91% (2.25)	99.18% (0.81)
	C	95.38% (4.00)	94.55% (1.00)	96.95% (0.48)
Adjusted Rand Index	N	0.9662 (0.07)	0.9677 (0.07)	0.9729 (0.0217)
	C	0.8706 (0.08)	0.8366 (0.04)	0.8538 (0.0453)

Several findings are concluded from Table 14. Under condition N, the BP method shows better performance in error and is found to be less sensitive than other methods with an error rate of 1.10% and a standard deviation of 0.02%. Further findings show that when condition C is imposed then Bernoulli has better performance in terms of findings. The ARI shows good measurements overall however the BP-Method under condition N has a very good ARI with a small standard deviation. The Average Purity of the BP-Method is the best out of all other methods, which is relevant to estimating coefficients accurately for the ZIP optimization.

6 Conclusion

In this paper, we extend the class of generalized linear mixture CWM models by accomplishing two main goals. First, we propose the methodology that allows for continuous covariates to follow a non-Gaussian distribution. Imposing Gaussian distribution on a skewed data may result in an suboptimal model fit. Second, we propose a new Poisson CWM methodology that uses Bernoulli-Poisson partitioning and allows for implementation of zero-inflated Poisson CWM model. We call our proposed model class GCWM a which reflects two extensions made to the existing CWM class of models.

The GCWM a models allows for great applications in predictive modeling of insurance claims by overcoming a few limitations of the current CWM models. Zero-inflated GCWM a allows for finding clusters within claims frequency which is an important information in risk classification and modeling of claims frequency. Further, some insurance rating variables used in the predictive modeling of severity claims may not strictly follow Gaussian assumptions, e.g. driver's age or car age (treated as continuous covariates). An adequate transformation can be considered on the continues covariates to relax current assumptions and improve the model fit. We demonstrated that if there is a need for transformation, a Lognormal transformation can be considered easily to improve the model fit.

The results of our extensive simulation study showed the excellent performance of the proposed model in case of modeling non-Gaussian covariates. We found that current CWM model fails to estimate the parameters accurately when the Gaussian assumption is violated. The GCWM a shows significant improvement in the model fit over the CWM model based on AIC and BIC criteria. We also tested Bernoulli-Poisson partitioning of zero-inflated GCWM a under different conditions and found that our proposed partitioning method has a very low misclassification rate, high average purity, and high average rand index.

Our approach is relevant to the actuarial pricing and risk management when current practices are based on implementation of various GLM models. Further extension of this work may incorporate Bayesian setting by exploring different assumption on informative and noninformative priors (see Ibrahim and Laud (1991)). By utilizing these techniques actuaries will be able to make inferences from posterior distributions of model parameters and from predictive distributions in order to improve pricing and risk management of the insured portfolio.

7 Appendix

7.1 Derivation of the Log-normal Distribution

Consider a random variable U having univariate log-normal distribution with parameters $\mu \in \mathbb{R}$ and $\sigma \in \mathbb{R}_+$. Have $u \in \mathbb{R}_+$, then the probability density function of random variable U is defined as [1]

$$\mathcal{LN}(u; \mu, \sigma) = \frac{1}{u\sigma\sqrt{2\pi}} \exp\left[-\frac{(\ln u - \mu)^2}{2\sigma^2}\right].$$

Further, if random variable X is normally distributed i.e. $X \sim \mathcal{N}(x; \mu, \sigma)$, then $U := \exp(X) \sim \mathcal{LN}(u; \mu, \sigma)$. To see this, let $p_U(u)$, and $p_X(x)$ be the probability density functions of U and X respectively. By the change of variables theorem (see Murphy and Bach (2012) section 2.6.2.1) the density $p_U(u)$ is derived as

$$p_U(u) = p_X(\ln u)\frac{\partial}{\partial u}\ln u = p_X(\ln u)\frac{1}{u} = \frac{1}{u\sigma\sqrt{2\pi}}\exp\left[-\frac{(\ln u - \mu)^2}{2\sigma^2}\right].$$

We extend to a log-normal multivariate case where the random variable $\boldsymbol{U}$ is parameterized by $\boldsymbol{\mu} \in \mathbb{R}^p$ and $\boldsymbol{\Sigma} \in \mathbb{R}_+{}^p$.

Lemma 7.1. *Let the random variable $\boldsymbol{X}$ have multivariate normal distribution ie. $\boldsymbol{X} \sim \mathcal{MVN}(\boldsymbol{x}, \boldsymbol{\mu}, \boldsymbol{\Sigma})$, then $\boldsymbol{U} := \exp(\boldsymbol{X}) \sim f^U(\boldsymbol{u}; \boldsymbol{\mu}, \boldsymbol{\Sigma})$. Here have $\boldsymbol{u} \in \mathbb{R}_+{}^p$ and the probability density function f^U is*

$$f^U(\boldsymbol{u}; \boldsymbol{\mu}, \boldsymbol{\Sigma}) = \frac{1}{(\prod_{i=1}^N u_i)|\boldsymbol{\Sigma}|(2\pi)^{\frac{p}{2}}}\exp\left[-\frac{1}{2}(\ln\boldsymbol{u} - \boldsymbol{\mu})'\boldsymbol{\Sigma}^{-1}(\ln\boldsymbol{u} - \boldsymbol{\mu})\right].$$

Proof. Let $f^U(\boldsymbol{u}; \boldsymbol{\mu}, \boldsymbol{\Sigma})$ and $f^X(\boldsymbol{x}; \boldsymbol{\mu}, \boldsymbol{\Sigma})$ be the probability density functions of $\boldsymbol{U}$ and $\boldsymbol{X}$ respectively. By the multivariate change of variables theorem (see Murphy and Bach (2012) section 2.6.2.1), we derive the log-normal distribution, where $|\det J_{\ln}(u)|$ is the absolute value of the determinant for the Jacobian of the multivariate transformation $\ln(\boldsymbol{U}) = \boldsymbol{X}$. Hence,

$$|\det J_{\ln}(\boldsymbol{u})| = \prod_{i=1}^n u_i^{-1}, \text{ and}$$

$$f^U(\boldsymbol{u}; \boldsymbol{\mu}, \boldsymbol{\Sigma}) = f^X(\ln\boldsymbol{u}; \boldsymbol{\mu}, \boldsymbol{\Sigma})|\det J_{\ln}(u)|$$

$$= f^X(\ln\boldsymbol{u}; \boldsymbol{\mu}, \boldsymbol{\Sigma})\prod_{i=1}^n u_i^{-1}$$

$$= \frac{1}{(\prod_{i=1}^N u_i)|\boldsymbol{\Sigma}|(2\pi)^{\frac{p}{2}}}\exp\left[-\frac{1}{2}(\ln\boldsymbol{u} - \boldsymbol{\mu})'\boldsymbol{\Sigma}^{-1}(\ln\boldsymbol{u} - \boldsymbol{\mu})\right].$$

$\square$

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[1] For full definition see Johnson et al. (1995)

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